

Amendments to the Specification:

Please replace the current Sequence Listing with the Sequence Listing attached hereto in computer readable format (CRF) and in paper format (**Exhibit 2**) (5 pages).

On page 6, please amend the paragraph at lines 19-21 as follows.

1A. TRT" [SEQ ID NO:1], a 244 bp fragment containing a direct repeat of the sequence AAAATCAGA (nucleotides 37-45 and 51-59 of SEQ ID NO:1), an inverted repeat of the sequence TAATACAACACAAT (nucleotides 111-124 of SEQ ID NO:1), and a direct repeat of the sequence ACGCAACACAATTTAT (nucleotides 175-190 and 211-226 of SEQ ID NO:1).

On page 6, please amend the paragraph at lines 22-24 as follows.

1B. A 134 bp fragment [SEQ ID NO:11] containing as the first 116 nucleotides TRT' [SEQ ID NO:2], a 116 bp fragment containing an inverted repeat of the sequence TAATACAACACAAT (nucleotides 1-14 of SEQ ID NO:2 and of SEQ ID NO:11) and a direct repeat of the sequence ACGCAACACAATTTAT (nucleotides 65-80 and 101-116 of SEQ ID NO:2 and of SEQ ID NO:11).

On page 6, please amend the paragraph at lines 25-26 as follows.

1C. TRT [SEQ ID NO:3], a 32 bp fragment containing an inverted repeat of the sequence TAATACAACACAAT (nucleotides 1-14 of SEQ ID NO:3).

On page 6, please amend the paragraph at lines 29-31 as follows.

2A ~~4A~~. A schematic diagram of the transposon Tn4430, with its 38 bp terminal inverted repeats (IR) represented by triangles and the position of the TnpI and TnpA (transposase) coding regions depicted as grey arrows.

On page 6, please amend the paragraph at lines 32-36 as follows.

2B ~~4B~~. Sequence of the 249 bp TRT¹ [SEQ ID NO:4] indicating the following: the Tn4430 left IR (first shaded box); TnpI binding regions (open boxes), within which is shown the inverted repeats (black arrows), and the conserved 9 bp sequence (5'-CAACACAAT-3') (nucleotides 118-126, 180-188, and 216-224 of SEQ ID NO:4) common to all four TnpI binding sites; the direct repeats (grey arrows); and the TnpI translational start site (second shaded box).